FIGURE 8

	Kinetics read at	wavelength 235 nm for a period of 2 minutes							
	12	27.506	NoFit	85.026	38.549	NoFit	-4.364	10.384	9.818
	=	-21.00 35.455 78.31 2 20.312 69.575 27	-4.255 301.55	NoFit	33.828 40.309 26.291 12.818 37.988 66.704 61.230 33.400 31.101 38.549	4.364	0.253	7.145 57.089 7.836 18.154 32.02	NoFit 1.165 3.018 46.200 0.000 9.823 99.536 396.76 9.818
	6	20.312	-4,255	134.56	33.400	NoFit	1.097	18.154	99.536
22	O	78.312	9,682	75.461 71.450 134.56	61.230	21.055	715.69 1.097	7,636	9.823
site 18	ထ	35.455	-2,240	75.461	66.704	NoFit		57,089	0000
utation	~	21.00	-2.786 -2.240	NoFit	37.988	5.922	1.968	7.145	46.200
s of mu	မ	NoFit	NOFIL	6.218	12.818		25.478 -7.968 4.155	7,636	3,019
l clone	2	380.25	NoFit	15,818	26.291	37.245 -16.36	8.385	31,309 7,636	1.165
V _{max} of GSSM clones of mutation site 182	4	10.691	59.177	20.911 341.11 45.818 6.218	40.309	NoFit	NoFit	5.544	NoFit
V _{max} of	0	NoFit	0.799	30,911	33.828	11.325	NoFit	-0.364	28.901
	2	48.338	.3313.539	26.455	24.278	3.173	13.046	4.149	
	_	34.031	357.33	2.787	0.853	16.869	10.766/13.046	0.476	9,684 51,590
The second secon		ৰ	0	ပ		u .	u.	<u>'</u> Ø	Ī